

## Sequence Listing

<120> Ar <130> Le <160> 2	Bayer Aktiengesellschaft Anti-Kazlauskas-Lipases LeA 35 991 2 PatentIn version 3.1												
<212> DN <213> ur <220> <221> CI	1 885 DNA unknown												
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Gln Leu I	atc gac tg Ile Asp Tr 35	g ccc gag p Pro Glu	gag ttc Glu Phe 40	atc cgg Ile Arg	ggg ctg Gly Leu 45	gct gaa Ala Glu	cga Arg	144					
ggc ttc c Gly Phe A 50	ogg gta at Arg Val Il	c tgt ttc e Cys Phe 55	gac aac Asp Asn	cgc gac Arg Asp	gcg ggg Ala Gly 60	ctt tcg Leu Ser	acg Thr	192					
	gaa ggc gt Glu Gly Va							240					
gcg agc a Ala Ser M	atg ggc ct Met Gly Le 85	a aag ccc u Lys Pro	agg gtg Arg Val	cct tac Pro Tyr 90	acc ctc Thr Leu	gac gac Asp Asp 95	atg Met	288					
	gac acc gt Asp Thr Va 100			Ala Leu				336					
His Val V	gtt ggc gt Val Gly Va 115							384					
	cac ggg ga His Gly Gl		Lys Ser					432					
tcc ggc a Ser Gly A 145	aac ccc cg Asn Pro Ar	c atg ccg g Met Pro 150	gcg ccc Ala Pro	agg ccg Arg Pro 155	cag gtg Gln Val	ctg caa Leu Gln	aag Lys 160	480					

								gat Asp									528
aac Asn	ttg Leu	gag Glu	ctt Leu 180	tta Leu	acc Thr	acc Thr	atc Ile	ggc Gly 185	agc Ser	ccc Pro	ggg ggg	ttg Leu	gac Asp 190	cgg Arg	gag Glu		576
								agc Ser									624
								gcc Ala									672
								gtc Val									720
								cag Gln									768
cat His	atc Ile	ccg Pro	gga Gly 260	gcc Ala	cgc Arg	ttc Phe	gag Glu	ctc Leu 265	atc Ile	gag Glu	ggc Gly	atg Met	ggg Gly 270	cac His	gac Asp		816
								ctg Leu									864
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<213> unknown

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Gly Phe Arg Val Ile Cys Phe Asp Asn Arg Asp Ala Gly Leu Ser Thr 50 55 60

Lys Leu Glu Gly Val Lys Lys Pro Asn Ile Ala Arg Val Phe Leu Leu 65 70 75 80

Ala Ser Met Gly Leu Lys Pro Arg Val Pro Tyr Thr Leu Asp Asp Met 85 90 95

Ala Leu Asp Thr Val Gly Leu Met Asp Ala Leu Gly Ile Glu Ser Thr 100 105 110

His Val Val Gly Val Ser Met Gly Gly Met Ile Ala Gln Ile Leu Gly 115 120 125

Ala Lys His Gly Glu Arg Val Lys Ser Leu Thr Leu Met Ile Thr Ser 130 135 140

Ser Gly Asn Pro Arg Met Pro Ala Pro Arg Pro Gln Val Leu Gln Lys 145 150 155 160

Phe Met Arg Val Pro Lys Ser Met Asp Lys Glu Glu Trp Ile Lys Tyr 165 170 175

Asn Leu Glu Leu Leu Thr Thr Ile Gly Ser Pro Gly Leu Asp Arg Glu 180 185 190

Lys Leu Ala Leu Asp Val Arg Lys Ser Ile Glu Arg Cys Leu Cys Pro 195 200 205

Glu Gly Thr Gln Arg Gln Leu Ala Ala Ile Leu Gln Ser Gly Ser Arg 210 215 220

Val Lys Leu Leu Arg Arg Ile Ala Val Pro Thr Leu Val Ile Ser Gly 225 230 235 240

Ala Glu Asp Pro Leu Leu Pro Tyr Gln Cys Gly Arg Asp Ile Ala Asp 245 250 255

His Ile Pro Gly Ala Arg Phe Glu Leu Ile Glu Gly Met Gly His Asp 260 265 270

Ile Pro Glu Arg His Ile Pro Arg Leu Ile Glu Leu Ile Ala Gly His 275 280 285

Ala Ala Ala Glu Ala 290